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## WHAT IS CLAIMED IS:

1. A method of inhibiting human telomerase activity comprising the step of contacting human telomerase with a polynucleotide comprising an antisense sequence of at least 7 nucleotides that specifically hybridizes to a nucleotide sequence within an accessible region of the RNA component of a human telomerase ("hTR"), but that does not hybridize to a sequence within a template region of the human telomerase, wherein the sequence within an accessible region is a sequence selected from nucleotides 137-193, 290-319, and 350-380 of hTR, whereby the polynucleotide inhibits the activity of the telomerase.

- 2. The method of claim 1 wherein the antisense sequence is between 10 and 50 nucleotides in length.
- 3. The method of claim 1 wherein the antisense sequence is between 15 and 35 nucleotides in length.
- 4. The method of claim 1 wherein the step of providing the cell with the polynucleotide comprises transfecting the cell with an expression vector comprising expression control sequences operatively linked to a nucleotide sequence encoding the antisense polynucleotide which vector expresses the polynucleotide.
  - 5. The method of claim 1 wherein the cell is a cancer cell.
- 6. A pharmaceutical composition comprising a pharmaceutically acceptable carrier and:
- (1) a polynucleotide comprising an antisense sequence of at least 7 nucleotides that specifically hybridizes to a nucleotide sequence within an accessible region of the RNA component of a human telomerase ("hTR"), but that does not hybridize to a sequence within a template region of the human telomerase, wherein the sequence within an accessible region is a sequence selected from nucleotides 137-193, 290-319, and 350-380 of hTR, or
- (2) an expression vector comprising expression control sequences operatively linked to a nucleotide sequence encoding the polynucleotide which vector expresses the polynucleotide.

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|----------------------------------|---|--|--|--|--|--|
|                                  | - 7. A method of treating a telomerase-related condition involving cells                    |  |  |  |  |  |
|                                  | exhibiting telomerase activity in a subject comprising the step of administering to the     |  |  |  |  |  |
|                                  | subject a pharmaceutical composition in an amount effective to inhibit telomerase activity  |  |  |  |  |  |
|                                  | in the cells, wherein the pharmaceutical composition comprises a pharmaceutically           |  |  |  |  |  |
|                                  | acceptable carrier and:   |  |  |  |  |  |
|                                  | (1) a polynucleotide comprising a sequence of at least 7 nucleotides                        |  |  |  |  |  |
|                                  | that specifically hybridizes to a nucleotide sequence within an accessible region of the    |  |  |  |  |  |
|                                  | RNA component of a human telomerase ("hTR"), but that does not hybridize to a               |  |  |  |  |  |
|                                  | sequence within a template region of the human telomerase, wherein the sequence within      |  |  |  |  |  |
|                                  | an accessible region is a sequence selected from nucleotides 137-193, 290-319, and 350-     |  |  |  |  |  |
|                                  | 380 of hTR, or  |  |  |  |  |  |
|                                  | (2) an expression vector comprising expression control sequences                            |  |  |  |  |  |
|                                  | operatively linked to a nucleotide sequence encoding the polynucleotide which vector        |  |  |  |  |  |
|                                  | expresses the antisense polynucleotide,   |  |  |  |  |  |
|                                  | whereby inhibiting telomerase activity in the cells provides the                            |  |  |  |  |  |
|                                  | treatment of the condition.   |  |  |  |  |  |
|                                  | ·   |  |  |  |  |  |
|                                  | 8. The method of claim 7 wherein the telomerase-related condition is                        |  |  |  |  |  |
|                                  | cancer and inhibition of telomerase activity in the cancer cells inhibits the growth of the |  |  |  |  |  |
|                                  | cancer.   |  |  |  |  |  |
|                                  |   |  |  |  |  |  |
|                                  | 9. The method of claim 7 wherein the pharmaceutical composition is                          |  |  |  |  |  |
|                                  | an injectable solution administered by injection.   |  |  |  |  |  |
|                                  |   |  |  |  |  |  |
|                                  | 10. The method of claim 7 wherein the pharmaceutical composition                            |  |  |  |  |  |
| comprises the polynucleotide     |   |  |  |  |  |  |
|                                  |   |  |  |  |  |  |
|                                  | 11. The method of claim 7 wherein the pharmaceutical composition                            |  |  |  |  |  |
| comprises the expression vector. |   |  |  |  |  |  |
|                                  |   |  |  |  |  |  |

A polynucleotide comprising an antisense sequence of at least 7

nucleotides that specifically hybridizes to a nucleotide sequence within an accessible

hybridize to a sequence within a template region of the human telomerase, wherein the

region of the RNA component of a human elomerase ("hTR"), but that does not

| 5  | sequence within an accessible region is a sequence selected from nucleotides 137-193,               |  |  |
|----|---|--|--|
| 6  | 290-319, and 350-380 of hTR.  |  |  |
| 1  | 13. The polynucleotide of claim 12 wherein the sequence is between 10                               |  |  |
| 2  | and 50 nucleotides in length.   |  |  |
| 1  | 14. The polynucleotide of claim 12 wherein the sequence is between 15                               |  |  |
| 2  | 14. The polynucleotide of claim 12 wherein the sequence is between 15 and 35 nucleotides in length. |  |  |
| 2  | and 33 nucleotides in length.   |  |  |
| 1  | 15. The polynucleotide of claim 12 whose sequence consists essentially                              |  |  |
| 2  | of the sequence within the an accessible region.  |  |  |
|    | •   |  |  |
| 1  | 16. The polynucleotide of claim 12 comprising DNA or RNA.   |  |  |
|    |   |  |  |
| 1  | 17. The polynucle of claim 12 comprising a nucleotide analog  |  |  |
| 2  | selected from phosphorothioates, phosphoramidates, methyl phosphonates, chiral-methyl               |  |  |
| 3. | phosphonates, 2-O-methyl ribonucleotides and peptide-nucleic acids.                                 |  |  |
|    |   |  |  |
| 1  | 18. The polynucled tide of claim 12 further comprising an inhibitory                                |  |  |
| 2  | moiety.   |  |  |
|    |   |  |  |
| 1  | 19. The polynucleo tide of claim 12 wherein the sequence is   |  |  |
| 2  | complementary to the nucleotide sequence within an accessible region.                               |  |  |
|    |   |  |  |
| 1  | 20. The polynucleotide of claim 12 which is at most 50 nucleotides                                  |  |  |
| 2  | long.   |  |  |
|    |   |  |  |
| I  | 21. The polynucleotide of claim 12 of less than about 50 nucleotides in                             |  |  |
| 2  | a sequence that specifically hybridizes to an accessible region of the RNA component of             |  |  |
| 3  | telomerase.   |  |  |
|    |   |  |  |
| 1  | 22. The polynucleotide of claim 12 whose nucleotide sequence is                                     |  |  |
| 2  | selected from the group consisting of   |  |  |
| 3  | CGT TCC TCT TCC TGC GGC CTG AAA CGG TGA (SEQ ID NO:2)   |  |  |

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| 4  |   | CGT TCC TCT TCC  | TGC GGC CT (SEQ ID NO:3)                |
|----|---|------------------|---|
| 5  | · | CGT TCC TCT TCC  | (SEQ ID NO:4)                           |
| 6  |   | CTG ACA GAG CC   | C AAC TCT TCG CGG TGG CAG (SEQ ID NO:5) |
| 7  |   | CTG ACA GAG CC   | AAO TCT TC (SEQ ID NO:6)                |
| 8  |   | CCA ACT CTT CGC  | GOT GGC AG (SEQ ID NO:7)                |
| 9  |   | GCT CTA GAA TGA  | ACG OTG GAA GGC GGC AGG (SEQ ID NO:8)   |
| 10 |   | GCT CTA GAA TG   | ACG GTG G (SEQ ID NO:9)                 |
| 11 |   | GCT CTA GAA TGA  | ACG (SEQ ID NO:10)                      |
| 12 |   | GCT CTA GAA TG   | (SEQ ID NO:11)                          |
| 13 |   | GCT CTA G (SEQ I | D NO:12)                                |
| 14 |   | CAT TIT TIG TIT  | GCT CTA GA (SEQ ID NO:13) and           |
| 15 |   | CGG GCC AGC AG   | C TGA CA (SEQ ID NO:14).                |
|    |   |                  | ,                                       |

- 23. An expression vector comprising a recombinant polynucleotide comprising expression control sequences operatively linked with a nucleotide sequence encoding a polynucleotide comprising an antisense sequence of at least 7 nucleotides that specifically hybridizes to a nucleotide sequence within an accessible region of the RNA component of a human telomerase ("hTR"), but that does not hybridize to a sequence within a template region of the human telorherase, wherein the sequence within an accessible region is a sequence selected from nucleotides 137-193, 290-319, and 350-380 of hTR.
- 24. The expression vector of claim 23 wherein the expression control sequences comprise a promoter selected from the metallothionein promoter, the constitutive adenovirus major late promoter, the dexamethasone-inducible MMTV promoter, the SV40 promoter, the MRP/polIII promoter, the constitutive MPSV promoter, the tetracycline-inducible CMV promoter (such as the human immediate-early CMV promoter), and the constitutive CMV promoter.
- 25. The expression vedtor of claim 23 wherein a viral vector or a plasmid vector comprising the recombinant polynucleotide.
- The expression vector of claim 25 wherein the vector is a plasmid 26. vector contained in a liposome.